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(xi)SEQU	JENCE DESCRIPTION: ST	BQ ID NO:1:				
Acucucuucc	GCAUCGCUGU	CUGCGAGGGC	CAGCUGUUGG	ecacecean	GAGGACAAAC	6 0
cuucocagu	CUUÜCCAGUA	CUCUUGGAUC	GGAAACCCGU	COOCCUCCOA	ACGUACUCCG	120
CCACCGAGGG	ACCUGAGCGA	GÜCCGCAÜCG	ACCGGAUCGG	AAAACCUCUC	GAGAAAGGCG	180
UČUAACCAGU	CACAGUCGCA		-		•	200
(2) INFORMATION					-	
	UENCE CHARACTERISTI (A) LENGTH: 33 base; (B) TYPE: mucleic scid (C) STRANDEDNESS: (D) TOPOLOGY: linear	aingle	:			
(ii)MOL	ECULE TYPE: mRNA				·	
(xi)SEQU	UENCE DESCRIPTION: S	EQ ID NO2:	, and the second second	•		
ACUCUCUUCC	acaucocuou	CUGCGAGGGC	CAG	•		3 3
(2) INFORMATION	FOR SEQ ID NO3:					
	UENCE CHARACTERIST (A) LENGTH: 12 base (B) TYPE: nucleic acid (C) STRANDEDNESS: (D) TOPOLOGY: linear	pairs single			,	
(ii) MOL	LECULE TYPE: DNA (Ben	omic)				-
(xi)SEQ	UENCE DESCRIPTION: S	SEQ ID NO3:	•			
AGCTTTGATC	AO -					1 2
(2) INFORMATION	FOR SEQ ID NO:4:	•				
(i)SEQ	UENCE CHARACTERIST (A) LENGTH: 12 base (B) TYPE: nucleic acid (C) STRANDEDNESS (D) TOPOLOGY: linea	pairs single			• ,	
(ii) MOI	LECULE TYPE: DNA (ge	nomic)		•		
(xi)SBQ	UENCE DESCRIPTION:	SEQ ID NO:4:				•
GCACCTGATC	· AA.			a to the state of	,	1 2
(2) INPORMATION	FOR SBQ ID NO:5:					
(i)SEQ	QUENCE CHARACTERIS: (A) LENGTH: 8 base; (B) TYPE: nucleic acid (C) STRANDEDNESS (D) TOPOLOGY: lines	pairs 1 I: single	•			
(ii) MO	LECULE TYPE: DNA (80	nomic)				
(xi)SEQ	QUENCE DESCRIPTION:	SBQ ID NO:5:	·			
GTGATCAA		•		,	,	8
(2) INFORMATION	N FOR SEQ ID NO:6:					

72

(i) SEQUENCE CHARACTERISTICS:
(A) LENOTH: 16 base pairs
(B) TYPE: mucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

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(ii)MO	LECULE TYPE: DNA (go:	omic)				
(xi)SEC	QUENCE DESCRIPTION:	SEQ ID NO:6:		•		
GATCTTGAT	CACTOCA	•				1
(2) INFORMATION	FOR SEQ ID NO:7:					
(i)sec	QUENCE CHARACTERIST (A) LENGTH: 8 base p (B) TYPE: nucleic acid (C) STRANDEDNESS: (D) TOPOLOGY: linear	airs : : single		,		
(ii)MO	LECULE TYPE: DNA (got	omic)	•	•		
(xi)SEQ	UENCE DESCRIPTION: S	SEQ ID NO:7:				
CGGATCCG				•		
(2) INPORMATION	FÓR SEQ ID NO:8:		•			
(i)seq	UENCE CHARACTERIST (A) LENGTH: 8 base p (B) TYPE: nucleic acid (C) STRANDEDNESS: (D) TOPOLOGY: linear	airs single			•	
(ii) MO	LECULE TYPE: DNA (gen	owic)				
(xi)SEQ	UENCE DESCRIPTION: S	EQ ID NO:8:				
CGGATCCG						8
(2) INFORMATION	FOR SEQ ID NO.9:					•
	UENCE CHARACTERIST (A) LENGTH: 287 base (B) TYPE: mucleic scid (C) STRANDEDNESS: (D) TOPOLOGY: linear ECULE TYPE: DNA (gen	pairs				
	UENCE DESCRIPTION: S	•				
		•	CGCTAGGGTG	CCGACGCGCA	TCTCGACTGC	6 0
ACGGT GCACC	AATGCTTCTG	GCGTCAGGCA	GCCAATCGGA	AGCTGTGGTA	TOGCTGTGCA	1 2 0
GGICGTATAĂ	TCACCGCATA	ATTCGAGTCG	CTCAAGGCGC	ACTCCCGTTC	CGGATAATGT	180
TTTTTOCTCC	GACATCATAA	CGGTTCCGGC	AAATATTCTG	AAATGAGCTG	TTGACAATTA	2 4 0
ATCATCGAAC	TAGTTAACTA	GTACGCAAGT	TCTCGTAAAA	AGGGTAT		287
(2) INFORMATION	FOR SEQ ID NO:10:		•	•		
• •	JENCE CHARACTERISTI (A) LENGTH: 285 base (B) TYPE: mucleic acid (C) STRANDEDNESS: (D) TOPOLOGY: linear	peirs single				
(ii)MOL	ECULE TYPE: DNA (geno	mic)				
(xi)SBQU	JENCE DESCRIPTION: SI	EQ ID NO:10:				
COATACCCTT	TTTACGAGAA	CTTGCGTACT	AGTTAACTAG	TTCGATGATT	AATTGTCAAC	60
OCTCATTTC	AGAATATTTG	CCGGAACCGT	TATGATGTCG	GAGCAAAAA	CATTATCCGG	120
ACGGGAGTG	CGCCTTGAGC	GACTCGAATT	ATGCGGTGAT	TATACGACCT	GCACAGCCAT	180
CCACAGCTT	CCGATTGGCT	GCCTGACGCC	AGAAGCATTG	GTGCACCGTG	CAGTCGAGAT	240

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GCGCGTCGGC ACCCTAGCGA CCACCGACCA TAACACCACA GCGTG	285
(2) INFORMATION FOR SEQ ID NO:11:	-
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 8 base pairs	
. (B) TYPE: nucleic acid	
(C) STRANDEDNESS: single (D) TOPOLOGY: linear	·
(i i) MOLECULE TYPE: DNA (genomic)	
(x i) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
CCATATGG	8
(2) INFORMATION FOR SEQ ID NO:12:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 8 base pairs	
(B) TYPE: moleic scid	
(C) STRANDEDNESS: single (D) TOPOLOGY: linear	·
(i i) MOLECULE TYPE: DNA (genomic)	
(x i) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
CCATATGG	8
A A DEPONALITION FOR ONE OF NO. 15.	
(2) INFORMATION FOR SEQ ID NO:13:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 8 base pairs	
(B) TYPE: mucleic acid	_
(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(i i) MOLECULE TYPE: DNA (genomic)	
(x i) SEQUENCE DESCRIPTION: SEQ ID NO:13:	•
CGTTAACG	
(2) Information for SEQ ID NO:14:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 8 base pairs	
(B) TYPE: micleic acid	
(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(i i) MOLECULE TYPE: DNA (genomio)	
(x i) SEQUENCE DESCRIPTION: SEQ ID NO:14:	
COTTAACG	8
(2) INFORMATION FOR SEQ ID NO:15:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 36 base pairs	-
(B) TYPE: modeio acid	
(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(i i) MOLECULE TYPE: DNA (genomic)	
(x i) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
QGGAAGTGCT GTGAAATATC CACCTGCGGC CTGAGA	3 6
(2) Information for seq ID No:16:	

-continued (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 46 base pairs (B) TYPE: modeic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:16: CTAGAGGGTA TTAATAATGT ATCGATTTAA ATAAGGAGGA ATAACA (2) INFORMATION FOR SEQ ID NO:17: (1) SEQUENCE CHARACTERISTICS: (A) LENGTH: 44 base pairs (B) TYPE: modeic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:17: TATGITATIC CICCITATIT AAATCGATAC ATTATTAATA CCCI (2) INFORMATION FOR SEQ ID NO:18: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: modeic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:18: GATCTATTAA CTCAATCTAG AC 22 (2) INFORMATION FOR SEQ ID NO:19: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:19: TCGAGTCTAG ATTGAGTTAA TA 2 2 (2) INFORMATION FOR SEQ ID NO:20: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 872 base pairs (B) TYPE: sucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:20: AAGCITTTCT CATTAAGGGA AGATTTCCCC AGGCAGCTCT TTCAAGGCCT AAAAGGTCCA 60 TGAGCTCCAT GGATTCTTCC CTGTTAAGAA CTTTATCCAT TTTTGCAAAA ATTGCAAAAG 120

AATAGGGATT TCCCCAAATA GTTTTGCTAG GCCTCAGAAA AAGCCTCCAC ACCCTTACTA

CTTGAGAGAA AGGGTGGAGG CAGAGGCGGC CTCGGCCTCT TATATATTAT AAAAAAAAAG

180

240

300



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GIGCATGACT	CACAGGGAA	TOCAGCCAAA	CCATGACCTC	AGGAAGGAAA	GTGCATGACT	360
CACAGGAGG	AGCTGCTTAC	CCATGGAATG	CAGCCAAACC	ATGACCTCAG	GAAGGAAAGT	420
GCATGACTGG	GCAGCCAGCC	AGTGGCAGTT	AATAGTGAAA	CCCCGCCGAC	AGACATGTTT	480
TGCGAGCCTA	GGAATCTTGG	CCTTGTCCCC	AGTTAAACTG	GACAAAGGCC	ATGGTTCTGC ·	5 4 0
GCCAGGCTGT	CCTTCGAGCG	втоттссосо	отсстсстсо	TATAGAAACT	CGGACCACTC	600
TGAGACGAAG	GCTCGCGTCC	AGGCCAGCAC	GAAGGAGGCT	AAGTGGGAGG	GGTAGCGGTC	660
GTTGTCCACT	AGGGGGTCCA	CTCGCTCCAG	GGTGTGAAGA	CACATOTCOC	CCTCTTCGGC	720
ATCAAGGAAG	GTGATTGGTT	TATAGGTĞTA	GGCCAGACCG	GOTOTTCCTG	AAGGGGGGCT	780
DDDDAAAAA	отооооосос	GTTCGTCCTC	ACTCTCTTCC	GCÀTCGCTGT	CTGCGAGGGC	8 4 0
CAGCTGATCA	GCCTAGGCTT	TGCAAAAAGC	TT			872

(2) INFORMATION FOR SEQ ID NO:21:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 643 base pairs
- (B) TYPE: mucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(i i) MOLECULE TYPE: DNA (genomic)

($\mathbf{x}(\mathbf{i})$) sequence description; seq id No:21:

AAGCTTTTCT	CATTAAGGGA	AGATTTCCCC	AGGCAGCTCT	TTCAAGGCCT	AAAAGGTCCA	, 60
TGAGCTCCAT	GGATTCTTCC	CTGTTAAGAA	CTTTATCCAT	TTTTGCAAAA	ATTGCAAAAG	1 2 0
TTADDDATAA	TCCCCAAATA	GTTTTGCTAG	GCCTCAGAAA	AAGCCTCCAC	ACCCTTACTA	180
CTTGAGAGAA	AGGGTGGAGG	CAGAGGCGGC	CTCGGCCTTC	TTATATATTA	TAAAAAAAA	2 4 0
GGCACÀGGG	AGGAGCTGCT	TACCCATGGA	ATGCAGCCAA	ACCATGACCT	CAGGAAGGAA	3 0 0
AGTGCATGAC	TCACAGGGGA	ATGCAGCCAA	ACCATGACCT	CAGGAAGGAA	AGTGCATGAC	360
TCACAGGGAG	GAGCTGCTTA	CCCATGGAAT	GCAGCCAAAC	CATGACCTCA	DAAADDAADD	4 2 0
TGCATGACTG	GGCAGCCAGC	CAGTGGCAGT	TAATACAGGG	TGTGAAGACA	CATGTCGCCC	480
TCTTCGGCAT	CAAGGAAGGT	GAATTGGTTT	ATAGGTGTAG	GCCACGTGAC	COOGTOTTCC	5 4 0
TOAAGGGGG	DDAAAATD	GGGTGGGGC	ocettcetcc	TCACTCTCTT	CCGCATCGCT	600
GTCTGCGAGG	GCCAGTGATC	AGCCTAGGCT	TTGCAAAAG	CTT		6 4 3

CM I 171,173

I claim:

- by inserting a vector comprising the DNA encoding human protein C into an adenovirus-transformed host cell then culturing said host cell under growth conditions suitable for production of said recombinant human protein C.
 - 2. The recombinant human protein C molecule of claim 1 wherein the adenovirus-transformed host cell is selected from the group consisting of AV12 cells and human embryonic kidney 293 cells.
 - 3. The recombinant human protein C molecule of claim 2 wherein the adenovirus-transformed host cell is an AV12 cell.
 - 4. The recombinant human protein C molecule of claim 2 wherein the adenovirus transformed host ceil is a human embryonic kidney 293 cell.